



IFW16

RAW SEQUENCE LISTING

DATE: 08/05/2004

PATENT APPLICATION: US/09/581,651C

TIME: 10:25:18

Input Set : A:\ERPO1.003APC.TXT

Output Set : N:\CRF4\08052004\I581651C.raw

4 <110> APPLICANT: Schor, Seth Lawrence
 5 Schor, Ana Maria
 7 <120> TITLE OF INVENTION: Polypeptides, Polynucleotides and Uses
 8 Thereof
 10 <130> FILE REFERENCE: ERPO1.003APC
 12 <140> CURRENT APPLICATION NUMBER: 09/581,651C
 13 <141> CURRENT FILING DATE: 2000-10-10
 15 <150> PRIOR APPLICATION NUMBER: PCT/GB98/03766
 16 <151> PRIOR FILING DATE: 1998-12-15
 18 <150> PRIOR APPLICATION NUMBER: GB 9726539.1
 19 <151> PRIOR FILING DATE: 1997-12-16
 21 <160> NUMBER OF SEQ ID NOS: 45
 23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 25 <210> SEQ ID NO: 1
 26 <211> LENGTH: 675
 27 <212> TYPE: PRT
 28 <213> ORGANISM: Homo sapiens
 30 <400> SEQUENCE: 1
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 32 1 5 10 15
 33 Leu Asn Met Leu Arg Gly Pro Gly Pro Gly Leu Leu Leu Ala Val
 34 20 25 30
 35 Leu Cys Leu Gly Thr Ala Val Pro Ser Thr Gly Ala Ser Lys Ser Lys
 36 35 40 45
 37 Arg Gln Ala Gln Gln Met Val Gln Pro Gln Ser Pro Val Ala Val Ser
 38 50 55 60
 39 Gln Ser Lys Pro Gly Cys Tyr Asp Asn Gly Lys His Tyr Gln Ile Asn
 40 65 70 75 80
 41 Gln Gln Trp Glu Arg Thr Tyr Leu Gly Asn Val Leu Val Cys Thr Cys
 42 85 90 95
 43 Tyr Gly Gly Ser Arg Gly Phe Asn Cys Glu Ser Lys Pro Glu Ala Glu
 44 100 105 110
 45 Glu Thr Cys Phe Asp Lys Tyr Thr Gly Asn Thr Tyr Arg Val Gly Asp
 46 115 120 125
 47 Thr Tyr Glu Arg Pro Lys Asp Ser Met Ile Trp Asp Cys Thr Cys Ile
 48 130 135 140
 49 Gly Ala Gly Arg Gly Arg Ile Ser Cys Thr Ile Ala Asn Arg Cys His
 50 145 150 155 160
 51 Glu Gly Gly Gln Ser Tyr Lys Ile Gly Asp Thr Trp Arg Arg Pro His
 52 165 170 175
 53 Glu Thr Gly Gly Tyr Met Leu Glu Cys Val Cys Leu Gly Asn Gly Lys
 54 180 185 190
 55 Gly Glu Trp Thr Cys Lys Pro Ile Ala Glu Lys Cys Phe Asp His Ala



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56          195          200          205
57 Ala Gly Thr Ser Tyr Val Val Gly Glu Thr Trp Glu Lys Pro Tyr Gln
58          210          215          220
59 Gly Trp Met Met Val Asp Cys Thr Cys Leu Gly Glu Gly Ser Gly Arg
60 225          230          235          240
61 Ile Thr Cys Thr Ser Arg Asn Arg Cys Asn Asp Gln Asp Thr Arg Thr
62          245          250          255
63 Ser Tyr Arg Ile Gly Asp Thr Trp Ser Lys Lys Asp Asn Arg Gly Asn
64          260          265          270
65 Leu Leu Gln Cys Ile Cys Thr Gly Asn Gly Arg Gly Glu Trp Lys Cys
66          275          280          285
67 Glu Arg His Thr Ser Val Gln Thr Thr Ser Ser Gly Ser Gly Pro Phe
68          290          295          300
69 Thr Asp Val Arg Ala Ala Val Tyr Gln Pro Gln Pro His Pro Gln Pro
70 305          310          315          320
71 Pro Pro Tyr Gly His Cys Val Thr Asp Ser Gly Val Val Tyr Ser Val
72          325          330          335
73 Gly Met Gln Trp Leu Lys Thr Gln Gly Asn Lys Gln Met Leu Cys Thr
74          340          345          350
75 Cys Leu Gly Asn Gly Val Ser Cys Gln Glu Thr Ala Val Thr Gln Thr
76          355          360          365
77 Tyr Gly Gly Asn Ser Asn Gly Glu Pro Cys Val Leu Pro Phe Thr Tyr
78          370          375          380
79 Asn Gly Arg Thr Phe Tyr Ser Cys Thr Thr Glu Gly Arg Gln Asp Gly
80 385          390          395          400
81 His Leu Trp Cys Ser Thr Thr Ser Asn Tyr Glu Gln Asp Gln Lys Tyr
82          405          410          415
83 Ser Phe Cys Thr Asp His Thr Val Leu Val Gln Thr Gln Gly Gly Asn
84          420          425          430
85 Ser Asn Gly Ala Leu Cys His Phe Pro Phe Leu Tyr Asn Asn His Asn
86          435          440          445
87 Tyr Thr Asp Cys Thr Ser Glu Gly Arg Arg Asp Asn Met Lys Trp Cys
88          450          455          460
89 Gly Thr Thr Gln Asn Tyr Asp Ala Asp Gln Lys Phe Gly Phe Cys Pro
90 465          470          475          480
91 Met Ala Ala His Glu Glu Ile Cys Thr Thr Asn Glu Gly Val Met Tyr
92          485          490          495
93 Arg Ile Gly Asp Gln Trp Asp Lys Gln His Asp Met Gly His Met Met
94          500          505          510
95 Arg Cys Thr Cys Val Gly Asn Gly Arg Gly Glu Trp Thr Cys Tyr Ala
96          515          520          525
97 Tyr Ser Gln Leu Arg Asp Gln Cys Ile Val Asp Asp Ile Thr Tyr Asn
98          530          535          540
99 Val Asn Asp Thr Phe His Lys Arg His Glu Glu Gly His Met Leu Asn
100 545          550          555          560
101 Cys Thr Cys Phe Gly Gln Gly Arg Gly Arg Trp Lys Cys Asp Pro Val
102          565          570          575
103 Asp Gln Cys Gln Asp Ser Glu Thr Gly Thr Phe Tyr Gln Ile Gly Asp
104          580          585          590

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105 Ser Trp Glu Lys Tyr Val His Gly Val Arg Tyr Gln Cys Tyr Cys Tyr
106      595      600      605
107 Gly Arg Gly Ile Gly Glu Trp His Cys Gln Pro Leu Gln Thr Tyr Pro
108      610      615      620
109 Ser Ser Ser Gly Pro Val Glu Val Phe Ile Thr Glu Thr Pro Ser Gln
110 625      630      635      640
111 Pro Asn Ser His Pro Ile Gln Trp Asn Ala Pro Gln Pro Ser His Ile
112      645      650      655
113 Ser Lys Tyr Ile Leu Arg Trp Arg Pro Val Ser Ile Pro Pro Arg Asn
114      660      665      670
115 Leu Gly Tyr
116      675
119 <210> SEQ ID NO: 2
120 <211> LENGTH: 2147
121 <212> TYPE: DNA
122 <213> ORGANISM: Homo sapiens
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126 ttaggggtcc ggggcccggg ctgctgctgc tggccgtcca gtgcctgggg acagcgggtgc 120
127 cctccacggg agcctcgaag agcaagaggc aggcctcagca aatgggttcag cccaggtccc 180
128 cgggtggtgt cagtcaaagc aagcccggtt gttatgacaa tggaaaacac tatcagataa 240
129 atcaacagtg ggagcggacc tacctaggca atgcgttggg ttgtacttgt tatggaggaa 300
130 gccgaggttt taactgcgag agtaaacctg aagctgaaga gacttgcttt gacaagtaca 360
131 ctgggaacac ttaccgagtg ggtgacactt atgagcgtcc taaagactcc atgatctggg 420
132 actgtacctg catcggggct gggcgaggga gaataagctg taccatcgca aaccgctgcc 480
133 atgaaggggg tcagtcctac aagattggtg acacctggag gagaccacat gagactgggtg 540
134 gttacatggt agagtgtgtg tgtcttggtg atggaaaagg agaatggacc tgcaagccca 600
135 tagctgagaa gtgttttgat catgctgctg ggacttccta tgtggtcgga gaaacgtggg 660
136 agaagcccta ccaaggctgg atgatggtag attgtacttg cctgggagaa ggcagcggac 720
137 gcatcacttg cacttctaga aatagatgca acgatcagga cacaaggaca tcctatagaa 780
138 ttggagacac ctggagcaag aaggataatc gaggaaacct gctccagtgc atctgcacag 840
139 gcaacggccg aggagagtgg aagtgtgaga ggcacacctc tgtgcagacc acatcgagcg 900
140 gatctggccc cttcaccgat gtcgtgcag ctgtttacca accgcagcct caccgccagc 960
141 ctccctcccta tggccactgt gtcacagaca gtggtgtggt ctactctgtg gggatgcagt 1020
142 ggctgaagac acaaggaaat aagcaaatgc tttgcacgtg cctgggcaac ggagtcagct 1080
143 gccaaagac agctgtaacc cagacttacg gtggcaactc aaatggagag ccatgtgtct 1140
144 taccattcac ctacaacgac aggacggaca gcacaacttc gaattatgag caggaccaga 1200
145 aatactcttt ctgcacagac cacactgttt tgggttcagac tcgaggagga aattccaatg 1260
146 gtgccttgtg ccacttcccc ttcctataca acaaccacaa ttacactgat tgcacttctg 1320
147 agggcagaag agacaacatg aagtgtgtgt ggaccacaca gaactatgat gccgaccaga 1380
148 agtttggtgt ctgccccatg gctgcccacg aggaaatctg cacaaccaat gaaggggtca 1440
149 tgtaccgcat tggagatcag tgggataagc agcatgacat gggtcacatg atgaggtgca 1500
150 cgtgtgttgg gaatggtcgt ggggaatgga catgcattgc ctactcgag ctctcgagatc 1560
151 agtgcattgt tgatgacatc acttacaatg tgaacgacac attccacaag cgtcatgaag 1620
152 aggggcacat gctgaactgt acatgcttcg gtcagggtcg gggcaggtgg aagtgtgatc 1680
153 ccgtcgacca atgccaggat tcagagactg ggacgtttta tcaaattgga gattcatggg 1740
154 agaagtatgt gcatggtgtc agataccagt gctactgcta tggccgtggc attggggagt 1800
155 ggcattgcc aaccttacag acctatccaa gctcaagtgg tcctgtcgaa gtatttatca 1860
156 ctgagactcc gagtgcagccc aactcccacc ccatccagtg gaatgcacca cagccatctc 1920

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157 acatttccaa gtacattctc aggtggagac ctgtgagtat cccacccaga aaccttggat 1980
158 actgagtctc ctaatcttat caattctgat ggtttctttt tttcccagct tttgagccaa 2040
159 caactctgat taactattcc tatagcattt actatatattg tttagtgaac aaacaatatg 2100
160 tgggtcaatta aattgacttg tagactgaaa aaaaaaaaaa aaaaaaa 2147
162 <210> SEQ ID NO: 3
163 <211> LENGTH: 20
164 <212> TYPE: PRT
165 <213> ORGANISM: Homo sapiens
167 <400> SEQUENCE: 3
168 Ile Ser Lys Tyr Ile Leu Arg Trp Arg Pro Val Ser Ile Pro Pro Arg
169 1 5 10 15
170 Asn Leu Gly Tyr
171 20
174 <210> SEQ ID NO: 4
175 <211> LENGTH: 21
176 <212> TYPE: PRT
177 <213> ORGANISM: Homo sapiens
179 <400> SEQUENCE: 4
180 Gln Gln Trp Glu Arg Thr Tyr Leu Gly Asn Ala Leu Val Cys Thr Cys
181 1 5 10 15
182 Tyr Gly Gly Ser Arg
183 20
186 <210> SEQ ID NO: 5
187 <211> LENGTH: 23
188 <212> TYPE: PRT
189 <213> ORGANISM: Homo sapiens
191 <400> SEQUENCE: 5
192 Pro Cys Val Leu Pro Phe Thr Tyr Asn Asp Arg Thr Asp Ser Thr Thr
193 1 5 10 15
194 Ser Asn Tyr Glu Gln Asp Gln
195 20
198 <210> SEQ ID NO: 6
199 <211> LENGTH: 20
200 <212> TYPE: PRT
201 <213> ORGANISM: Homo sapiens
203 <400> SEQUENCE: 6
204 Thr Asp His Thr Val Leu Val Gln Thr Arg Gly Gly Asn Ser Asn Gly
205 1 5 10 15
206 Ala Leu Cys His
207 20
210 <210> SEQ ID NO: 7
211 <211> LENGTH: 21
212 <212> TYPE: PRT
213 <213> ORGANISM: Homo sapiens
215 <400> SEQUENCE: 7
216 Val Gly Asn Gly Arg Gly Glu Trp Thr Cys Ile Ala Tyr Ser Gln Leu
217 1 5 10 15
218 Arg Asp Gln Cys Ile
219 20

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Input Set : A:\ERPO1.003APC.TXT

Output Set: N:\CRF4\08052004\I581651C.raw

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222 <210> SEQ ID NO: 8
223 <211> LENGTH: 21
224 <212> TYPE: PRT
225 <213> ORGANISM: Homo sapiens
227 <400> SEQUENCE: 8
228 Gln Gln Trp Glu Arg Thr Tyr Leu Gly Asn Val Leu Val Cys Thr Cys
229 1 5 10 15
230 Tyr Gly Gly Ser Arg
231 20
234 <210> SEQ ID NO: 9
235 <211> LENGTH: 39
236 <212> TYPE: PRT
237 <213> ORGANISM: Homo sapiens
239 <400> SEQUENCE: 9
240 Glu Pro Cys Val Leu Pro Phe Thr Tyr Asn Gly Arg Thr Phe Tyr Ser
241 1 5 10 15
242 Cys Thr Thr Thr Glu Arg Gln Asp Gly His Leu Trp Cys Ser Thr Thr
243 20 25 30
244 Ser Asn Tyr Glu Gln Asp Gln
245 35
248 <210> SEQ ID NO: 10
249 <211> LENGTH: 21
250 <212> TYPE: PRT
251 <213> ORGANISM: Homo sapiens
253 <400> SEQUENCE: 10
254 Cys Thr Asp His Thr Val Leu Val Gln Thr Gln Gly Gly Asn Ser Asn
255 1 5 10 15
256 Gly Ala Leu Cys His
257 20
260 <210> SEQ ID NO: 11
261 <211> LENGTH: 21
262 <212> TYPE: PRT
263 <213> ORGANISM: Homo sapiens
265 <400> SEQUENCE: 11
266 Val Gly Asn Gly Arg Gly Glu Trp Thr Cys Thr Ala Tyr Ser Gln Leu
267 1 5 10 15
268 Arg Asp Gln Cys Ile
269 20
272 <210> SEQ ID NO: 12
273 <211> LENGTH: 20
274 <212> TYPE: PRT
275 <213> ORGANISM: Homo sapiens
277 <400> SEQUENCE: 12
278 Ile Ser Lys Thr Ile Leu Arg Trp Arg Pro Lys Asn Ser Val Gly Arg
279 1 5 10 15
280 Trp Lys Glu Ala
281 20
284 <210> SEQ ID NO: 13
285 <211> LENGTH: 18

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/05/2004
PATENT APPLICATION: US/09/581,651C TIME: 10:25:19

Input Set : A:\ERP01.003APC.TXT
Output Set: N:\CRF4\08052004\I581651C.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:37; Xaa Pos. 676,679,683,717

VERIFICATION SUMMARY

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Input Set : **A:\ERPO1.003APC.TXT**

Output Set: **N:\CRF4\08052004\I581651C.raw**

L:763 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:672

L:767 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:704